Figure la

CAG Gln 1	GTG Val	CAG Gln	CTG Leu	GTG Val 5	GAG Glu	TCT Ser	GGG Gly	GGA Gly	GGC Gly 10	GTT Val	GTG Val	CAG Gln	CCT Pro	GGA Gly IS	AGG Arg	48
TCC Ser	CTG Leu	AGA Arg	CTC Leu 20	TCC Ser	TGT:	GCA `Ala	GCC Ala	TCT Ser 25	GGA Gly	TTC Phe	ACC Thr	TTC Phe	AGT Ser 30	AGC Ser	TAT Tyr	96
GAC Asp	ATG Met	TCT Ser 35	TGG Trp	GTT Val	CGC Arg	CAG Gln	GCT Ala 40	CCG Pro	GGC Gly	AAG Lys	GGT Gly	CTG Leu 45	GAG Glu	TGS TIP	GTC Val	144
GCA Ala	AAA Lys 50	GTT Val	AGT Ser	AGT Ser	GGT Gly	GGT Gly 55	GGT Gly	AGC Ser	ACC Thr	TAC Tyr	TAT Tyr 60	TTA Leu	GAĆ Asp	ACT Thr	GTG Val	192
CAG Gln 65	GGC Gly	CGA A r g	TTC Phe	ACC Thr	ATC Ile 70	TCC Ser	AGA Arg	GAC Asp	TAA neA	AGT Ser 75	AAG Lys	DAA neA	ACC Thr	CTA Leu	TAC Tyr 80	240
CTG Leu	CAA Gln	ATG Met	AAC Asn	TCT Ser 85	CTG Leu	AGA Arg	GCC Ala	GAG Glu	GAC Asp 90	aca Thr	GCC Ala	GTG Val	TAT Tyr	TAC TYT 95	TGT Cys	288
GCA Ala	AGA Arg	CAT Eis	AAC Asn 100	TAC Tyr	GLY GGC	AGT Ser	TTT Phe	GCT Ala 105	TAC TYI	TGG Trp	GGC Gly	CAA Gln	GGG Gly 110	ACT Thr	aca Thr	336
GTG Val	ACT Thr	GTT Val 115	TCT Ser	AGT Ser												351

Figure 1b

+ 3				ACC Thr 10					
95				AGC ser					
144				CAA Gln					
192	GGC Gly			ATC Ile				arg	
243				ACT Thr					
288				CAG Gln 90					
321				ATT Ile			Gly		

Figure 2a

GAA Glu l	GTG Val	CAG Gln	CTG Leu	GTG Val 5	GAG Glu	TCT Ser	GGG Gly	GGA Gly	GGC Gly 10	TTA Leu	GTG Val	AAG Lys	CCT Pro	GGA Gly 13	agg Arg	48
TCC Ser	CTG Leu	AGA Arg	CTC Leu 20	TCC Ser	TGT Cys	GCA Ala	GCC Ala	TCT Ser 25	GGA Gly	TTC Phe	GCT Ala	TTC Phe	AGT Ser 30	AGC Se:	TAT Tyr	9 6
GAC Asp	ATG Met	TCT Ser 35	TGG Trp	GTT Val	CGC Arg	cag Gln	ATT Ile 40	CCG Pro	GAG Glu	AAG Lys	AGG Arg	CTG Leu 45	GAG Glu	TGG TT	GTC Val	144
GCA Ala	AAA Lys 50	GTT Val	AGT Ser	AGT Ser	GGT Gly	GGT Gly 55	GGT Gly	AGC Ser	ACC Thr	TAC Tyr	TAT TYT 60	TTA Leu	GAC Asp	ACT Thr	GTG Val	192
CAG Gln 65	GGC Gly	CGA Arg	TTC Phe	ACC Thr	ATC Ile 70	TCC Ser	AGA Arg	GAC Asp	AAT Asn	GCC Ala 75	AAG Lys	AAC Asn	ACC Thr	CTA Leu	TAC TYT 80	240
CTG Leu	CAA Gln	ATG Het	AGC Ser	AGT Ser 85	CTG Leu	AAC Asn	TCT Ser	GAG Glu	GAC Asp 90	ACA Thr	GCC Ala	ATG Met	TAT Tyr	TAC Ty: 93	TGT Cys	288
GCA Ala	AGA Arg	CAT Bis	AAC Asn 100	TAC Tyr	GGC	AGT Ser	TTT Phe	GCT Ala 105	TAC Tyr	TGG Trp	GLY	CAA Gln	GGG Gly 110	ACT The	CTG Leu	336
	ACT Thr		Ser													351

Figure 2b

	l ATT													48	
	AGC Ser													9 6	
	A CAC 1 His													144	
	G TAT B TYT 50	λIg												192	:
	r gga r gly s													240	;
	A GAT u Asp				Tyr					ser			His	288	3
AC Th	G TTC r Phe	GGA Gly	GLY	Gly	Thr	Lys	Leu	GAA Glu	Ile	AAG Lys				32	1

LM609 Competition Assay

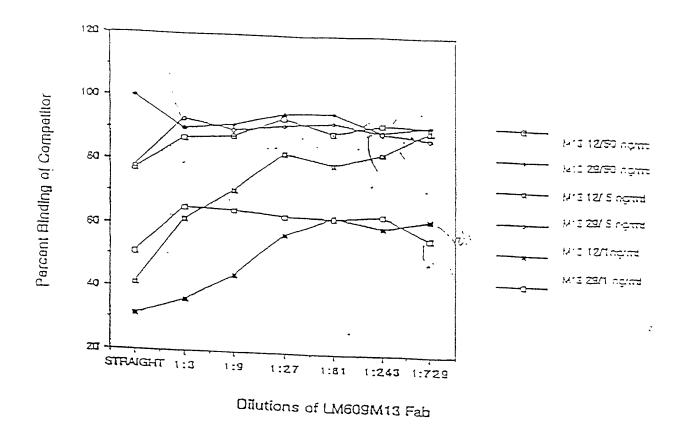
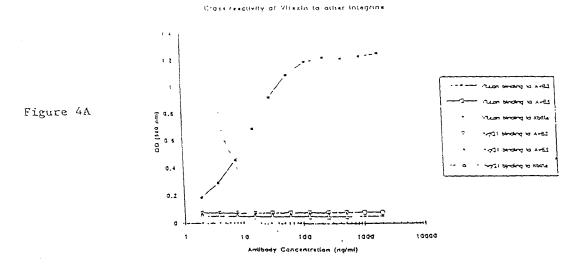


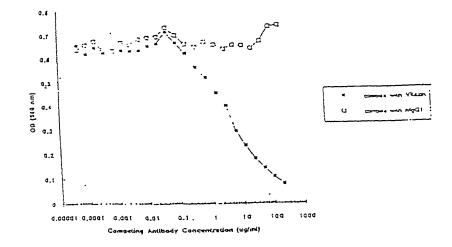
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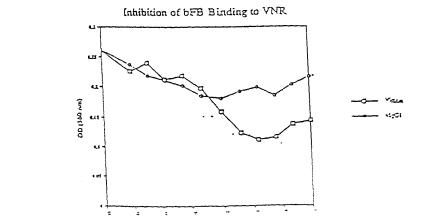
Figure 4B

Figure 4C

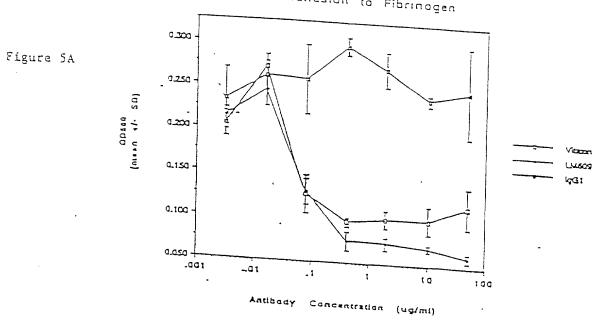




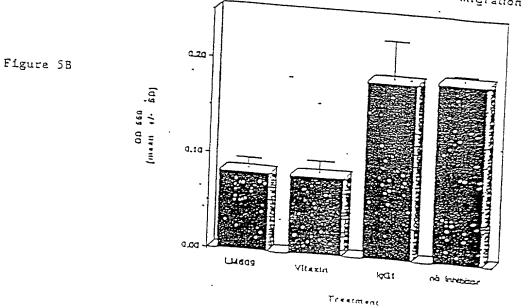


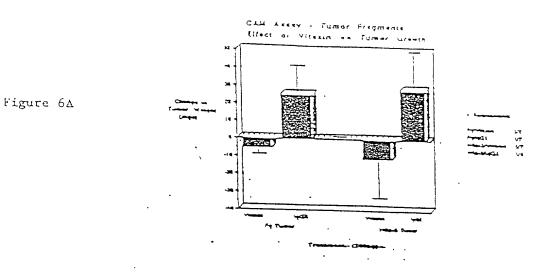


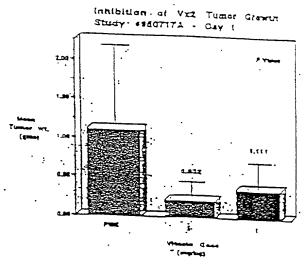
Inhibition of M21 Melanoma Cell Adhesion to Fibrinogen



Inhibition of Endothelial Cell Migration







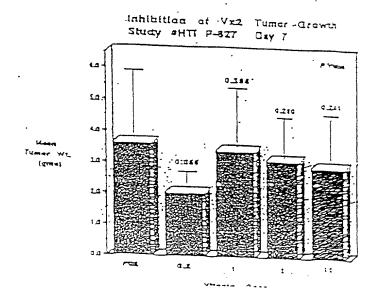


Figure 6C

Figure 6B

Figure 7

							CTC Leu			48
							ATT Ile			96
							AGG AFG 45			144
							GCC Ala 60			192
							AGT Ser			240
			Tyr			Ser	AGC Ser		His	288
		Gly			ATT					321

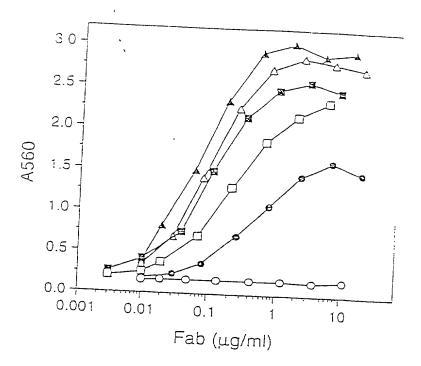


FIGURE 8

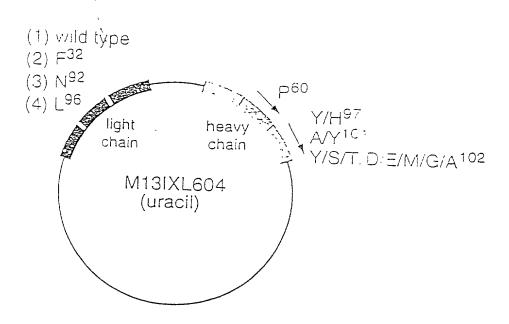
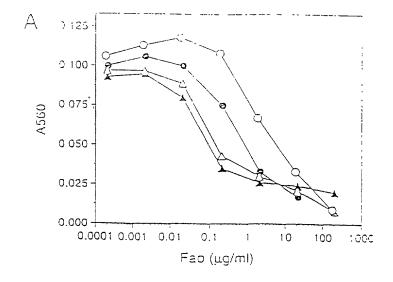


FIGURE 9



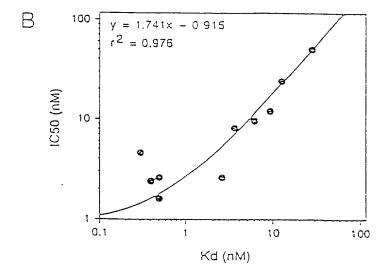


FIGURE 10

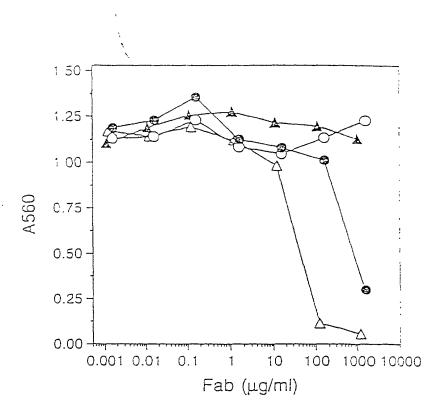


FIGURE 11